

## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

## (i) APPLICANT:

- (A) NAME: AKZO NOBEL N.V.  
(B) STREET: Velperweg 76  
(C) CITY: Arnhem  
(E) COUNTRY: The Netherlands  
(F) POSTAL CODE (ZIP): 6824 BM

(ii) TITLE OF INVENTION: Oligonucleotides for the amplification and detection of Epstein Barr Virus (EBV) nucleic acid

(iii) NUMBER OF SEQUENCES: 36

## (iv) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

## (2) INFORMATION FOR SEQ ID NO: 1:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 23 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GCCGGTGTGT TGTTTCGTATA TGG

23

## (2) INFORMATION FOR SEQ ID NO: 2:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

CTCCCTTTAC AACCTAAGGC

20

## (2) INFORMATION FOR SEQ ID NO: 3:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 25 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

AGAGACAAGG TCCTTAATCG CATCC

25

## (2) INFORMATION FOR SEQ ID NO: 4:

WO 99/45155

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

AATACAGACA ATGGACTCCC

20

## (2) INFORMATION FOR SEQ ID NO: 5:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

CGTCTCCCCT TTGGAATGGC CCCTGGACCC

30

## (2) INFORMATION FOR SEQ ID NO: 6:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

CGGGCGGACC AGCTGTACTT GA

22

(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 23 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

GAGGTTTTGA TAGGGAGAGG AGA

23

(2) INFORMATION FOR SEQ ID NO: 8:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 22 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

09623329 11.1300

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

22

(2) INFORMATION FOR SEQ ID NO: 9:

(D) TOPOLOGY: linear

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

22

(2) INFORMATION FOR SEQ ID NO: 10:

(D) TOPOLOGY: linear

(vi) ORIGINAL SOURCE:

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

19

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

21

(2) INFORMATION FOR SEQ ID NO: 12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs  
(B) TYPE: nucleic acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

ATACCTAAGA CAAGTTTGCT

20

(2) INFORMATION FOR SEQ ID NO: 13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

ATCAACCAAT AGAGTCCACC A

21

(2) INFORMATION FOR SEQ ID NO: 14:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

0962329 4100

CATCGTTATG AGTGACTGGA

20

## (2) INFORMATION FOR SEQ ID NO: 15:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 23 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

ACTGATGATC ACCCTCCTGC TCA

23

## (2) INFORMATION FOR SEQ ID NO: 16:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA to mRNA

## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

GGACAGGCAT TGTCCTTGG

20

## (2) INFORMATION FOR SEQ ID NO: 17:

00623329-11300



(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

TAACTGTGGT TTCCATGACG

20

(2) INFORMATION FOR SEQ ID NO: 18:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:

AGGTACTCTT GGTGCAGCCC

20

(2) INFORMATION FOR SEQ ID NO: 19:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs

09623320 11300

WO 99/45155

- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:

AGCATATAGG AACAGTCGTG CC

22

(2) INFORMATION FOR SEQ ID NO: 20:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

AGTGGACATG AAGAGCACGA A

21

(2) INFORMATION FOR SEQ ID NO: 21:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

00623339-11300

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

AGCTCTGGCA CTGCTAGCGT CACTGATTTT

30

(2) INFORMATION FOR SEQ ID NO: 22:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:

CAGGTTTCATC GCTCAGCTCC

20

(2) INFORMATION FOR SEQ ID NO: 23:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

00623320.1.1.300

- (vi) ORIGINAL SOURCE:
  - (A) ORGANISM: Epstein-Barr virus

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

GGCTGTCACC GCTTTCTTGG

20

- (2) INFORMATION FOR SEQ ID NO: 24:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 19 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: cDNA to mRNA

- (vi) ORIGINAL SOURCE:
  - (A) ORGANISM: Epstein-Barr virus

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

AGTGTTGGCA CTTCTGTGG

19

- (2) INFORMATION FOR SEQ ID NO: 25:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 21 base pairs
  - (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: cDNA to mRNA

- (vi) ORIGINAL SOURCE:
  - (A) ORGANISM: Epstein-Barr virus

09623329.11300

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:

AGCATGGGAG ATGTTGGCAG C

21

(2) INFORMATION FOR SEQ ID NO: 26:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:

CTGGTTTAAA CTGGGCCCCAG GAGAGGAGCA

30

(2) INFORMATION FOR SEQ ID NO: 27:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

WO 99/45155

20

TGGAGCGAAG GTTAGTGGTC

(2) INFORMATION FOR SEQ ID NO: 28:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

22

TACCTGGCAC CTGAGTGTGG AG

(2) INFORMATION FOR SEQ ID NO: 29:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

- (A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

24

AGAATTGGAT CATTCTGAC AGGG

25

(A) ORGANISM: Epstein-Barr virus

30

(i) SEQUENCE CHARACTERISTICS:

Geographical location		Study period		Study design		Study population		Study results	
Country	Region	Start	End	Design	Sample size	Age range	Gender	Prevalence	Incidence
USA	California	1990	1992	Cross-sectional	1,000	18-74	Male	1.2%	0.5%
USA	Florida	1990	1992	Cross-sectional	1,000	18-74	Female	1.5%	0.6%
USA	Illinois	1990	1992	Cross-sectional	1,000	18-74	Male	1.8%	0.7%
USA	Michigan	1990	1992	Cross-sectional	1,000	18-74	Female	2.1%	0.8%
USA	New York	1990	1992	Cross-sectional	1,000	18-74	Male	2.4%	0.9%
USA	Ohio	1990	1992	Cross-sectional	1,000	18-74	Female	2.7%	1.0%
USA	Pennsylvania	1990	1992	Cross-sectional	1,000	18-74	Male	3.0%	1.1%
USA	Texas	1990	1992	Cross-sectional	1,000	18-74	Female	3.3%	1.2%
USA	Virginia	1990	1992	Cross-sectional	1,000	18-74	Male	3.6%	1.3%
USA	Washington	1990	1992	Cross-sectional	1,000	18-74	Female	3.9%	1.4%
USA	Wisconsin	1990	1992	Cross-sectional	1,000	18-74	Male	4.2%	1.5%
USA	Zoo	1990	1992	Cross-sectional	1,000	18-74	Female	4.5%	1.6%
USA	Illinois	1990	1992	Cross-sectional	1,000	18-74	Male	4.8%	1.7%
USA	Michigan	1990	1992	Cross-sectional	1,000	18-74	Female	5.1%	1.8%
USA	New York	1990	1992	Cross-sectional	1,000	18-74	Male	5.4%	1.9%
USA	Ohio	1990	1992	Cross-sectional	1,000	18-74	Female	5.7%	2.0%
USA	Pennsylvania	1990	1992	Cross-sectional	1,000	18-74	Male	6.0%	2.1%
USA	Texas	1990	1992	Cross-sectional	1,000	18-74	Female	6.3%	2.2%
USA	Virginia	1990	1992	Cross-sectional	1,000	18-74	Male	6.6%	2.3%
USA	Washington	1990	1992	Cross-sectional	1,000	18-74	Female	6.9%	2.4%
USA	Wisconsin	1990	1992	Cross-sectional	1,000	18-74	Male	7.2%	2.5%
USA	Zoo	1990	1992	Cross-sectional	1,000	18-74	Female	7.5%	2.6%
USA	Illinois	1990	1992	Cross-sectional	1,000	18-74	Male	7.8%	2.7%
USA	Michigan	1990	1992	Cross-sectional	1,000	18-74	Female	8.1%	2.8%
USA	New York	1990	1992	Cross-sectional	1,000	18-74	Male	8.4%	2.9%
USA	Ohio	1990	1992	Cross-sectional	1,000	18-74	Female	8.7%	3.0%
USA	Pennsylvania	1990	1992	Cross-sectional	1,000	18-74	Male	9.0%	3.1%
USA	Texas	1990	1992	Cross-sectional	1,000	18-74	Female	9.3%	3.2%
USA	Virginia	1990	1992	Cross-sectional	1,000	18-74	Male	9.6%	3.3%
USA	Washington	1990	1992	Cross-sectional	1,000	18-74	Female	9.9%	3.4%
USA	Wisconsin	1990	1992	Cross-sectional	1,000	18-74	Male	10.2%	3.5%
USA	Zoo	1990	1992	Cross-sectional	1,000	18-74	Female	10.5%	3.6%
USA	Illinois	1990	1992	Cross-sectional	1,000	18-74	Male	10.8%	3.7%
USA	Michigan	1990	1992	Cross-sectional	1,000	18-74	Female	11.1%	3.8%
USA	New York	1990	1992	Cross-sectional	1,000	18-74	Male	11.4%	3.9%
USA	Ohio	1990	1992	Cross-sectional	1,000	18-74	Female	11.7%	4.0%
USA	Pennsylvania	1990	1992	Cross-sectional	1,000	18-74	Male	12.0%	4.1%
USA	Texas	1990	1992	Cross-sectional	1,000	18-74	Female	12.3%	4.2%
USA	Virginia	1990	1992	Cross-sectional	1,000	18-74	Male	12.6%	4.3%
USA	Washington	1990	1992	Cross-sectional	1,000	18-74	Female	12.9%	4.4%
USA	Wisconsin	1990	1992	Cross-sectional	1,000	18-74	Male		

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

CTACCTTCCA CGACTTCACC

20

(2) INFORMATION FOR SEQ ID NO: 33:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

AAGTCTTTTA TAAGGCTCCG GC

22

(2) INFORMATION FOR SEQ ID NO: 34:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single



(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

AGGCCATGGT GTCATCCATC

20

(2) INFORMATION FOR SEQ ID NO: 35:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(vi) ORIGINAL SOURCE:

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

AGAGAGAGAG TAGGTCCGCG G

21

(2) INFORMATION FOR SEQ ID NO: 36:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 30 base pairs

(B) TYPE: nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA to mRNA

(A) ORGANISM: Epstein-Barr virus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

30

Descriptive statistics		Pearson correlation		Spearman correlation		Kendall correlation	
Variable	Mean	Variable	Mean	Variable	Mean	Variable	Mean
1. Age	35.5	2. Age	35.5	3. Age	35.5	4. Age	35.5
5. Age	35.5	6. Age	35.5	7. Age	35.5	8. Age	35.5
9. Age	35.5	10. Age	35.5	11. Age	35.5	12. Age	35.5
13. Age	35.5	14. Age	35.5	15. Age	35.5	16. Age	35.5
17. Age	35.5	18. Age	35.5	19. Age	35.5	20. Age	35.5
21. Age	35.5	22. Age	35.5	23. Age	35.5	24. Age	35.5
25. Age	35.5	26. Age	35.5	27. Age	35.5	28. Age	35.5
29. Age	35.5	30. Age	35.5	31. Age	35.5	32. Age	35.5
33. Age	35.5	34. Age	35.5	35. Age	35.5	36. Age	35.5
37. Age	35.5	38. Age	35.5	39. Age	35.5	40. Age	35.5
41. Age	35.5	42. Age	35.5	43. Age	35.5	44. Age	35.5
45. Age	35.5	46. Age	35.5	47. Age	35.5	48. Age	35.5
49. Age	35.5	50. Age	35.5	51. Age	35.5	52. Age	35.5
53. Age	35.5	54. Age	35.5	55. Age	35.5	56. Age	35.5
57. Age	35.5	58. Age	35.5	59. Age	35.5	60. Age	35.5
61. Age	35.5	62. Age	35.5	63. Age	35.5	64. Age	35.5
65. Age	35.5	66. Age	35.5	67. Age	35.5	68. Age	35.5
69. Age	35.5	70. Age	35.5	71. Age	35.5	72. Age	35.5
73. Age	35.5	74. Age	35.5	75. Age	35.5	76. Age	35.5
77. Age	35.5	78. Age	35.5	79. Age	35.5	80. Age	35.5
81. Age	35.5	82. Age	35.5	83. Age	35.5	84. Age	35.5
85. Age	35.5	86. Age	35.5	87. Age	35.5	88. Age	35.5
89. Age	35.5	90. Age	35.5	91. Age	35.5	92. Age	35.5
93. Age	35.5	94. Age	35.5	95. Age	35.5	96. Age	35.5
97. Age	35.5	98. Age	35.5	99. Age	35.5	100. Age	35.5